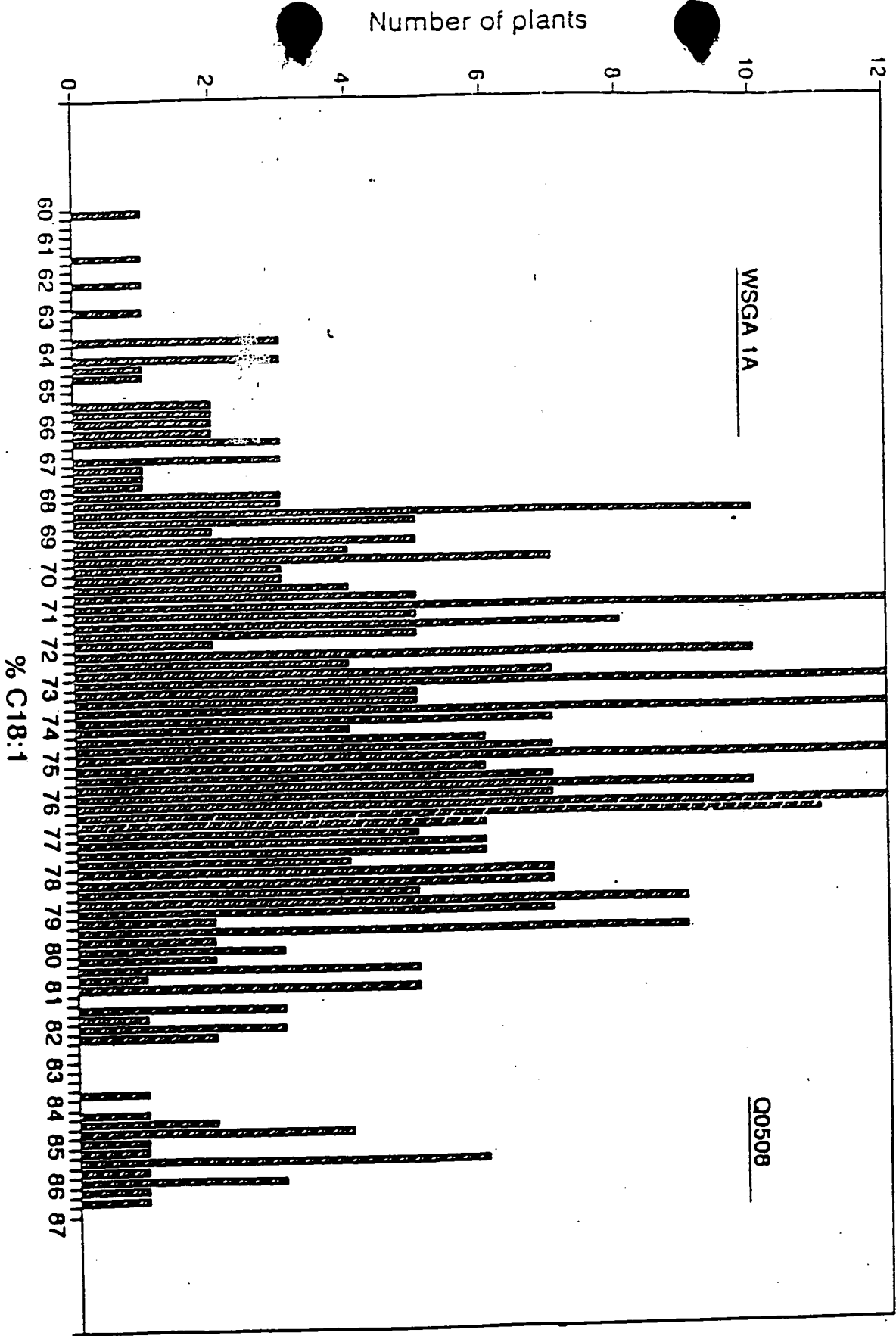


Fig. 1 C18:1 Frequencies
for 92EF (MSGA 1A X Q0508)



0971904.012901

		10	20	30	40	
1	ATGGG	TGCAG	GTGG	AAGA	ATGCA	AGTGTCTCCTCTCTCCA Fad2-D wt
1	ATGGG	TGCAG	GTGG	AAGA	ATGCA	AGTGTCTCCTCTCTCCA Fad2-D (GA316) IMC129
1	ATGGG	TGCAG	GTGG	AAGA	ATGCA	AGTGTCTCCTCTCTCCA Fad2-F wt
1	ATGGG	TGCAG	GTGG	AAGA	ATGCA	AGTGTCTCCTCTCTCCA Fad2-F (TA515) Q508
1	ATGGG	TGCAG	GTGG	AAGA	ATGCA	AGTGTCTCCTCTCTCCA Fad2-F (GA908) Q4275
		50	60	70	80	
41	AAAAG	TCTGA	AAACCG	ACAA	ACATCA	AGCGCGGTACCCCTGCGGA Fad2-D wt
41	AAAAG	TCTGA	AAACCG	ACAA	ACATCA	AGCGCGGTACCCCTGCGGA Fad2-D (GA316) IMC129
41	AGAAG	TCTGA	AAACCG	ACAA	ACATCA	AGCGCGGTACCCCTGCGGA Fad2-F wt
41	AGAAG	TCTGA	AAACCG	ACAA	ACATCA	AGCGCGGTACCCCTGCGGA Fad2-F (TA515) Q508
41	AGAAG	TCTGA	AAACCG	ACAA	ACATCA	AGCGCGGTACCCCTGCGGA Fad2-F (GA908) Q4275
		90	100	110	120	
81	GACAC	CGCCCT	TTCA	CTGT	CGGAG	AACTCAAGAAAGCAATC Fad2-D wt
81	GACAC	CGCCCT	TTCA	CTGT	CGGAG	AACTCAAGAAAGCAATC Fad2-D (GA316) IMC129
81	GACAC	CGCCCT	TTCA	CTGT	CGGAG	AACTCAAGAAAGCAATC Fad2-F wt
81	GACAC	CGCCCT	TTCA	CTGT	CGGAG	AACTCAAGAAAGCAATC Fad2-F (TA515) Q508
81	GACAC	CGCCCT	TTCA	CTGT	CGGAG	AACTCAAGAAAGCAATC Fad2-F (GA908) Q4275
		130	140	150	160	
121	CCACC	GCACT	GTTT	CAAAC	GGCTCG	ATCCCTCGCTCTTTCT Fad2-D wt
121	CCACC	GCACT	GTTT	CAAAC	GGCTCG	ATCCCTCGCTCTTTCT Fad2-D (GA316) IMC129
121	CCACC	GCACT	GTTT	CAAAC	GGCTCG	ATCCCTCGCTCTTTCT Fad2-F wt
121	CCACC	GCACT	GTTT	CAAAC	GGCTCG	ATCCCTCGCTCTTTCT Fad2-F (TA515) Q508
121	CCACC	GCACT	GTTT	CAAAC	GGCTCG	ATCCCTCGCTCTTTCT Fad2-F (GA908) Q4275
		170	180	190	200	
161	CCTAC	CTCAT	CTGGG	GACAT	CATCAT	ATAGCCTCCTGCTTCTA Fad2-D wt
161	CCTAC	CTCAT	CTGGG	GACAT	CATCAT	ATAGCCTCCTGCTTCTA Fad2-D (GA316) IMC129
161	CCTAC	CTCAT	CTGGG	GACAT	CATCAT	ATAGCCTCCTGCTTCTA Fad2-F wt
161	CCTAC	CTCAT	CTGGG	GACAT	CATCAT	ATAGCCTCCTGCTTCTA Fad2-F (TA515) Q508
161	CCTAC	CTCAT	CTGGG	GACAT	CATCAT	ATAGCCTCCTGCTTCTA Fad2-F (GA908) Q4275
		210	220	230	240	
201	CTACG	TGCGC	CACTT	ACTT	CCCTCT	CCTCCCTCACCCT Fad2-D wt
201	CTACG	TGCGC	CACTT	ACTT	CCCTCT	CCTCCCTCACCCT Fad2-D (GA316) IMC129
201	CTACG	TGCGC	CACTT	ACTT	CCCTCT	CCTCCCTCACCCT Fad2-F wt
201	CTACG	TGCGC	CACTT	ACTT	CCCTCT	CCTCCCTCACCCT Fad2-F (TA515) Q508
201	CTACG	TGCGC	CACTT	ACTT	CCCTCT	CCTCCCTCACCCT Fad2-F (GA908) Q4275
		250	260	270	280	
241	CTCTC	CCTAC	TTG	CGCCT	GCGCT	CTCTACTGGGGCCTGCCAAG Fad2-D wt
241	CTCTC	CCTAC	TTG	CGCCT	GCGCT	CTCTACTGGGGCCTGCCAAG Fad2-D (GA316) IMC129
241	CTCTC	CCTAC	TTG	CGCCT	GCGCT	CTCTACTGGGGCCTGCCAAG Fad2-F wt
241	CTCTC	CCTAC	TTG	CGCCT	GCGCT	CTCTACTGGGGCCTGCCAAG Fad2-F (TA515) Q508
241	CTCTC	CCTAC	TTG	CGCCT	GCGCT	CTCTACTGGGGCCTGCCAAG Fad2-F (GA908) Q4275

FIG. 2A

	290	300	310	320	
281	GCTGCGTCCTAACC	GGCGTCTGGGTCATAGCCCA	CACTG	Fad2-D wt	
281	GCTGCGTCCTAACC	GGCGTCTGGGTCATAGCCCA	CAAGTG	Fad2-D (GA316)	IMC129
281	GGTGCCTCCTAACC	GGCGTCTGGGTCATAGCCCA	CAAGTG	Fad2-F wt	
281	GGTGCCTCCTAACC	GGCGTCTGGGTCATAGCCCA	CAAGTG	Fad2-F (TA515)	Q508
281	GGTGCCTCCTAACC	GGCGTCTGGGTCATAGCCCA	CAAGTG	Fad2-F (GA908)	Q4275
	330	340	350	360	
321	CGGCCACCAACGCCCTTCAGCG	ACTACCAAGTGGCTGGACGAC	Fad2-D wt		
321	CGGCCACCAACGCCCTTCAGCG	ACTACCAAGTGGCTGGACGAC	Fad2-D (GA316)	IMC129	
321	CGGCCACCAACGCCCTTCAGCG	ACTACCAAGTGGCTGGACGAC	Fad2-F wt		
321	CGGCCACCAACGCCCTTCAGCG	ACTACCAAGTGGCTGGACGAC	Fad2-F (TA515)	Q508	
321	CGGCCACCAACGCCCTTCAGCG	ACTACCAAGTGGCTGGACGAC	Fad2-F (GA908)	Q4275	
	370	380	390	400	
361	ACCGTCGGGCTCATCTTCC	ACTCCTTCCTCCTCGTCCCTT	Fad2-D wt		
361	ACCGTCGGGCTCATCTTCC	ACTCCTTCCTCCTCGTCCCTT	Fad2-D (GA316)	IMC129	
361	ACCGTCGGGCTCATCTTCC	ACTCCTTCCTCCTCGTCCCTT	Fad2-F wt		
361	ACCGTCGGGCTCATCTTCC	ACTCCTTCCTCCTCGTCCCTT	Fad2-F (TA515)	Q508	
361	ACCGTCGGGCTCATCTTCC	ACTCCTTCCTCCTCGTCCCTT	Fad2-F (GA908)	Q4275	
	410	420	430	440	
401	ACTTCTCCTGGAAGTACAGT	CATCGACGCCACCATTC	CAA	Fad2-D wt	
401	ACTTCTCCTGGAAGTACAGT	CATCGACGCCACCATTC	CAA	Fad2-D (GA316)	IMC129
401	ACTTCTCCTGGAAGTACAGT	CATCGACGCCACCATTC	CAA	Fad2-F wt	
401	ACTTCTCCTGGAAGTACAGT	CATCGACGCCACCATTC	CAA	Fad2-F (TA515)	Q508
401	ACTTCTCCTGGAAGTACAGT	CATCGACGCCACCATTC	CAA	Fad2-F (GA908)	Q4275
	450	460	470	480	
441	CACTGGCTCCCTCGAGAGAG	ACGAAGTGTTTGTCCCCAAG	Fad2-D wt		
441	CACTGGCTCCCTCGAGAGAG	ACGAAGTGTTTGTCCCCAAG	Fad2-D (GA316)	IMC129	
441	CACTGGCTCCCTCGAGAGAG	ACGAAGTGTTTGTCCCCAAG	Fad2-F wt		
441	CACTGGCTCCCTCGAGAGAG	ACGAAGTGTTTGTCCCCAAG	Fad2-F (TA515)	Q508	
441	CACTGGCTCCCTCGAGAGAG	ACGAAGTGTTTGTCCCCAAG	Fad2-F (GA908)	Q4275	
	490	500	510	520	
481	AAGAAGTTCAGACATCAAGT	GGTACGGCAAGTACCTCAACA	Fad2-D wt		
481	AAGAAGTTCAGACATCAAGT	GGTACGGCAAGTACCTCAACA	Fad2-D (GA316)	IMC129	
481	AAGAAGTTCAGACATCAAGT	GGTACGGCAAGTACCTCAACA	Fad2-F wt		
481	AAGAAGTTCAGACATCAAGT	GGTACGGCAAGTACCTCAACA	Fad2-F (TA515)	Q508	
481	AAGAAGTTCAGACATCAAGT	GGTACGGCAAGTACCTCAACA	Fad2-F (GA908)	Q4275	
	530	540	550	560	
521	ACCCTTTTGGGACGCACC	GTGATGTTAACGGTTTCAGTT	CAC	Fad2-D wt	
521	ACCCTTTTGGGACGCACC	GTGATGTTAACGGTTTCAGTT	CAC	Fad2-D (GA316)	IMC129
521	ACCCTTTTGGGACGCACC	GTGATGTTAACGGTTTCAGTT	CAC	Fad2-F wt	
521	ACCCTTTTGGGACGCACC	GTGATGTTAACGGTTTCAGTT	CAC	Fad2-F (TA515)	Q508
521	ACCCTTTTGGGACGCACC	GTGATGTTAACGGTTTCAGTT	CAC	Fad2-F (GA908)	Q4275

FIG. 2B

	570	580	590	600	
561	TCTCGGCTGGCCCTTTGTACTTAGCCCTTCAACGTC	CGGGG	Fad2-D wt		
561	TCTCGGCTGGCCCTTTGTACTTAGCCCTTCAACCTCTCGGGG		Fad2-D (GA316) IMC129		
561	TCTCGGCTGGCCGTTGTACTTAGCCCTTCAACGTCCTCGGGGA		Fad2-F wt		
561	TCTCGGCTGGCCGTTGTACTTAGCCCTTCAACCTCTCGGGGA		Fad2-F (TA515) Q508		
561	TCTCGGCTGGCCGTTGTACTTAGCCCTTCAACGTCCTCGGGGA		Fad2-F (GA908) Q4275		
	610	620	630	640	
601	AGACCTTACGACGGCGGGCTTTCGCTTGGCCATTTTCCACCCCCA		Fad2-D wt		
601	AGACCTTACGACGGCGGGCTTTCGCTTGGCCATTTTCCACCCCCA		Fad2-D (GA316) IMC129		
601	AGACCTTACGACGGCGGGCTTTCGCTTGGCCATTTTCCACCCCCA		Fad2-F wt		
601	AGACCTTACGACGGCGGGCTTTCGCTTGGCCATTTTCCACCCCCA		Fad2-F (TA515) Q508		
601	AGACCTTACGACGGCGGGCTTTCGCTTGGCCATTTTCCACCCCCA		Fad2-F (GA908) Q4275		
	650	660	670	680	
641	ACGCTCCCCATCTACAACGACCGGTGACCGTCTCCAGATATA		Fad2-D wt		
641	ACGCTCCCCATCTACAACGACCGGTGACCGTCTCCAGATATA		Fad2-D (GA316) IMC129		
641	ACGCTCCCCATCTACAACGACCGCGGACCGTCTCCAGATATA		Fad2-F wt		
641	ACGCTCCCCATCTACAACGACCGCGGACCGTCTCCAGATATA		Fad2-F (TA515) Q508		
641	ACGCTCCCCATCTACAACGACCGCGGACCGTCTCCAGATATA		Fad2-F (GA908) Q4275		
	690	700	710	720	
681	CATCTCCGACGCTGGGCATCCTCGCCGTCCTGCTACGGGTCTC		Fad2-D wt		
681	CATCTCCGACGCTGGGCATCCTCGCCGTCCTGCTACGGGTCTC		Fad2-D (GA316) IMC129		
681	CATCTCCGACGCTGGGCATCCTCGCCGTCCTGCTACGGGTCTC		Fad2-F wt		
681	CATCTCCGACGCTGGGCATCCTCGCCGTCCTGCTACGGGTCTC		Fad2-F (TA515) Q508		
681	CATCTCCGACGCTGGGCATCCTCGCCGTCCTGCTACGGGTCTC		Fad2-F (GA908) Q4275		
	730	740	750	760	
721	TACCGCTACGCTGCTGTCCAAAGGAGTTGCCCTCGATGGGTCT		Fad2-D wt		
721	TACCGCTACGCTGCTGTCCAAAGGAGTTGCCCTCGATGGGTCT		Fad2-D (GA316) IMC129		
721	TTCCGTTACGCCGCCGCCGAGGGAGTGGCCCTCGATGGGTCT		Fad2-F wt		
721	TTCCGTTACGCCGCCGCCGAGGGAGTGGCCCTCGATGGGTCT		Fad2-F (TA515) Q508		
721	TTCCGTTACGCCGCCGCCGAGGGAGTGGCCCTCGATGGGTCT		Fad2-F (GA908) Q4275		
	770	780	790	800	
761	GCTTCTACGGAGTTCCCTCTTCTGATTGTCAACGGGGTTCTT		Fad2-D wt		
761	GCTTCTACGGAGTTCCCTCTTCTGATTGTCAACGGGGTTCTT		Fad2-D (GA316) IMC129		
761	GCTTCTACGGAGTCCCGCTTCTGATTGTCAATGGGTTTCTT		Fad2-F wt		
761	GCTTCTACGGAGTCCCGCTTCTGATTGTCAATGGGTTTCTT		Fad2-F (TA515) Q508		
761	GCTTCTACGGAGTCCCGCTTCTGATTGTCAATGGGTTTCTT		Fad2-F (GA908) Q4275		
	810	820	830	840	
801	AGTTTTTGATCACTTACTTTGCAGCACACGCATCCTTCCCTG		Fad2-D wt		
801	AGTTTTTGATCACTTACTTTGCAGCACACGCATCCTTCCCTG		Fad2-D (GA316) IMC129		
801	CGTGTTGATCACTTACTTTGCAGCACACGCATCCTTCCCTG		Fad2-F wt		
801	CGTGTTGATCACTTACTTTGCAGCACACGCATCCTTCCCTG		Fad2-F (TA515) Q508		
801	CGTGTTGATCACTTACTTTGCAGCACACGCATCCTTCCCTG		Fad2-F (GA908) Q4275		

FIG. 2C

	850	860	870	880	
841	CCTCACTATGACTCTCTCTGAGTGGGATTGGTTGAGGGGAG				Fad2-D wt
841	CCTCACTATGACTCTCTCTGAGTGGGATTGGTTGAGGGGAG				Fad2-D (GA316) IMC129
841	CCTCACTACGATTCTCTCCGAGTGGGATTGGTTGAGGGGAG				Fad2-F wt
841	CCTCACTACGATTCTCTCCGAGTGGGATTGGTTGAGGGGAG				Fad2-F (TA515) Q508
841	CCTCACTACGATTCTCTCCGAGTGGGATTGGTTGAGGGGAG				Fad2-F (GA908) Q4275
	890	900	910	920	
881	CTTTGGCCACCGTTGACACAGACTACCGAATCTTGAACAA				Fad2-D wt
881	CTTTGGCCACCGTTGACACAGACTACCGAATCTTGAACAA				Fad2-D (GA316) IMC129
881	CTTTGGCTACCGTTGACACAGACTACCGAATCTTGAACAA				Fad2-F wt
881	CTTTGGCTACCGTTGACACAGACTACCGAATCTTGAACAA				Fad2-F (TA515) Q508
881	CTTTGGCTACCGTTGACACAGACTACCGAATCTTGAACAA				Fad2-F (GA908) Q4275
	930	940	950	960	
921	GGTCTTCCACAATATCACGGACACGCACGTGGCGGCATCAC				Fad2-D wt
921	GGTCTTCCACAATATCACGGACACGCACGTGGCGGCATCAC				Fad2-D (GA316) IMC129
921	GGTCTTCCACAATATTACCGACACGCACGTGGCGGCATCAT				Fad2-F wt
921	GGTCTTCCACAATATTACCGACACGCACGTGGCGGCATCAT				Fad2-F (TA515) Q508
921	GGTCTTCCACAATATTACCGACACGCACGTGGCGGCATCAT				Fad2-F (GA908) Q4275
	970	980	990	1000	
961	CTGTTCTTCGACCATGCCCGCATTATCATGCGATGGAAGCTA				Fad2-D wt
961	CTGTTCTTCGACCATGCCCGCATTATCATGCGATGGAAGCTA				Fad2-D (GA316) IMC129
961	CTGTTCTTCACGATGCCCGCATTATCACGCGATGGAAGCTA				Fad2-F wt
961	CTGTTCTTCACGATGCCCGCATTATCACGCGATGGAAGCTA				Fad2-F (TA515) Q508
961	CTGTTCTTCACGATGCCCGCATTATCACGCGATGGAAGCTA				Fad2-F (GA908) Q4275
	1010	1020	1030	1040	
1001	CGAAGGCGGATAAAAGCCGATACTGGGAGAGTATTATCAGTT				Fad2-D wt
1001	CGAAGGCGGATAAAAGCCGATACTGGGAGAGTATTATCAGTT				Fad2-D (GA316) IMC129
1001	CCAAGGCGGATAAAAGCCGATACTGGGAGAGTATTATCAGTT				Fad2-F wt
1001	CCAAGGCGGATAAAAGCCGATACTGGGAGAGTATTATCAGTT				Fad2-F (TA515) Q508
1001	CCAAGGCGGATAAAAGCCGATACTGGGAGAGTATTATCAGTT				Fad2-F (GA908) Q4275
	1050	1060	1070	1080	
1041	CGATGGGACGCGCGGTGGTTAAGGCGATGTGGAGGGGAGGCG				Fad2-D wt
1041	CGATGGGACGCGCGGTGGTTAAGGCGATGTGGAGGGGAGGCG				Fad2-D (GA316) IMC129
1041	CGATGGGACGCGCGGTGGTTAAGGCGATGTGGAGGGGAGGCG				Fad2-F wt
1041	CGATGGGACGCGCGGTGGTTAAGGCGATGTGGAGGGGAGGCG				Fad2-F (TA515) Q508
1041	CGATGGGACGCGCGGTGGTTAAGGCGATGTGGAGGGGAGGCG				Fad2-F (GA908) Q4275
	1090	1100	1110	1120	
1081	AAGGAGTGATCTATGTGGAACCGGACAGGCCAAGGTGAGA				Fad2-D wt
1081	AAGGAGTGATCTATGTGGAACCGGACAGGCCAAGGTGAGA				Fad2-D (GA316) IMC129
1081	AAGCAGTGATCTATGTGGAACCGGACAGGCCAAGGTGAGA				Fad2-F wt
1081	AAGGAGTGATCTATGTGGAACCGGACAGGCCAAGGTGAGA				Fad2-F (TA515) Q508
1081	AAGCAGTGATCTATGTGGAACCGGACAGGCCAAGGTGAGA				Fad2-F (GA908) Q4275

FIG. 2D

1130 1140 1150
 1121 AGAAAGGTGTGTTCTGGTACAACAATAAGTTATGA
 1121 AGAAAGGTGTGTTCTGGTACAACAATAAGTTATGA
 1121 AGAAAGGTGTGTTCTGGTACAACAATAAGTTATGA
 1121 AGAAAGGTGTGTTCTGGTACAACAATAAGTTATGA
 1121 AGAAAGGTGTGTTCTGGTACAACAATAAGTTATGA

Pad2-D wt
 Pad2-D (GA316) TMC129
 Pad2-F wt
 Pad2-P (TA515) Q508
 Pad2-F (GA908) Q4275

FIG. 2E

0974904-01904
 T062T0-106T260

		10	20	
1	Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser Glu Thr Asp Asn			Fad2-D wt
1	Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser Glu Thr Asp Asn			Fad2-D (GA316) IMC129
1	Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser Glu Thr Asp Thr			Fad2-F wt
1	Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser Glu Thr Asp Thr			Fad2-F (TA515) Q508
1	Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser Glu Thr Asp Thr			Fad2-F (GA908) Q4275
		30	40	
61	Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile			Fad2-D wt
61	Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile			Fad2-D (GA316) IMC129
61	Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile			Fad2-F wt
61	Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile			Fad2-F (TA515) Q508
61	Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile			Fad2-F (GA908) Q4275
		50	60	
121	Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile			Fad2-D wt
121	Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile			Fad2-D (GA316) IMC129
121	Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile			Fad2-F wt
121	Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile			Fad2-F (TA515) Q508
121	Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile			Fad2-F (GA908) Q4275
		70	80	
181	Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			Fad2-D wt
181	Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			Fad2-D (GA316) IMC129
181	Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			Fad2-F wt
181	Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			Fad2-F (TA515) Q508
181	Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			Fad2-F (GA908) Q4275
		90	100	
241	Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val			Fad2-D wt
241	Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val			Fad2-D (GA316) IMC129
241	Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val			Fad2-F wt
241	Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val			Fad2-F (TA515) Q508
241	Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val			Fad2-F (GA908) Q4275
		110	120	
301	Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp			Fad2-D wt
301	Trp Val Ile Ala His Lys Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp			Fad2-D (GA316) IMC129
301	Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp			Fad2-F wt
301	Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp			Fad2-F (TA515) Q508
301	Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp			Fad2-F (GA908) Q4275
		130	140	
361	Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser			Fad2-D wt
361	Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser			Fad2-D (GA316) IMC129
361	Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser			Fad2-F wt
361	Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser			Fad2-F (TA515) Q508
361	Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser			Fad2-F (GA908) Q4275

FIG. 3A

	150	160	
421	His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys	Fad2-D wt	
421	His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys	Fad2-D (GA316)	IMC129
421	His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys	Fad2-F wt	
421	His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys	Fad2-F (TA515)	Q508
421	His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys	Fad2-F (GA908)	Q4275
	170	180	
481	Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val	Fad2-D wt	
481	Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val	Fad2-D (GA316)	IMC129
481	Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val	Fad2-F wt	
481	Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr His Asn Asn Pro Leu Gly Arg Thr Val	Fad2-F (TA515)	Q508
481	Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val	Fad2-F (GA908)	Q4275
	190	200	
541	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Fad2-D wt	
541	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Fad2-D (GA316)	IMC129
541	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Fad2-F wt	
541	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Fad2-F (TA515)	Q508
541	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Fad2-F (GA908)	Q4275
	210	220	
601	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Fad2-D wt	
601	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Fad2-D (GA316)	IMC129
601	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Fad2-F wt	
601	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Fad2-F (TA515)	Q508
601	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Fad2-F (GA908)	Q4275
	230	240	
661	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu	Fad2-D wt	
661	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu	Fad2-D (GA316)	IMC129
661	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu	Fad2-F wt	
661	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu	Fad2-F (TA515)	Q508
661	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu	Fad2-F (GA908)	Q4275
	250	260	
721	Tyr Arg Tyr Ala Ala Val Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu	Fad2-D wt	
721	Tyr Arg Tyr Ala Ala Val Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu	Fad2-D (GA316)	IMC129
721	Phe Arg Tyr Ala Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu	Fad2-F wt	
721	Phe Arg Tyr Ala Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu	Fad2-F (TA515)	Q508
721	Phe Arg Tyr Ala Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu	Fad2-F (GA908)	Q4275
	270	280	
781	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu	Fad2-D wt	
781	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu	Fad2-D (GA316)	IMC129
781	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu	Fad2-F wt	
781	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu	Fad2-F (TA515)	Q508
781	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu	Fad2-F (GA908)	Q4275

FIG. 3B

	290	300	
841	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg	Fad2-D wt	
841	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg	Fad2-D (GA316) IMC129	
841	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg	Fad2-F wt	
841	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg	Fad2-F (TA515) Q508	
841	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg	Fad2-F (GA908) Q4275	
	310	320	
901	Asp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His	Fad2-D wt	
901	Asp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His	Fad2-D (GA316) IMC129	
901	Asp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His	Fad2-F wt	
901	Asp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His	Fad2-F (TA515) Q508	
901	Asp Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His	Fad2-F (GA908) Q4275	
	330	340	
961	Leu Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Fad2-D wt	
961	Leu Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Fad2-D (GA316) IMC129	
961	Leu Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Fad2-F wt	
961	Leu Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Fad2-F (TA515) Q508	
961	Leu Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Fad2-F (GA908) Q4275	
	350	360	
1021	Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Fad2-D wt	
1021	Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Fad2-D (GA316) IMC129	
1021	Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Fad2-F wt	
1021	Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Fad2-F (TA515) Q508	
1021	Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Fad2-F (GA908) Q4275	
	370	380	
1081	Lys Glu Cys Ile Tyr Val Glu Pro Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	Fad2-D wt	
1081	Lys Glu Cys Ile Tyr Val Glu Pro Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	Fad2-D (GA316) IMC129	
1081	Lys Glu Cys Ile Tyr Val Glu Pro Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	Fad2-F wt	
1081	Lys Glu Cys Ile Tyr Val Glu Pro Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	Fad2-F (TA515) Q508	
1081	Lys Glu Cys Ile Tyr Val Glu Pro Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	Fad2-F (GA908) Q4275	
1141	Asn Asn Lys Leu ter	Fad2-D wt	
1141	Asn Asn Lys Leu ter	Fad2-D (GA316) IMC129	
1141	Asn Asn Lys Leu ter	Fad2-F wt	
1141	Asn Asn Lys Leu ter	Fad2-F (TA515) Q508	
1141	Asn Asn Lys Leu ter	Fad2-F (GA908) Q4275	

FIG. 3C